Figure 1

Alignment Of Conserved Amino Acid Sequences

| | AKPPIIVTRDSPNALRSHVLEVSPGADIVESVSTYARRGRGVSVLGGNGTVSNVTLRQ PKPPTIITRDSPNVLRSHVLEVTSGSDISEAVSTYATRRGCGVCIISGTGAVTNVTIRQ | VVTLHGRFEILSLTGTVLPPPAPPGAGGLSIFLA VITLHGRFDILSLTGTALPPPAPPGAGGLTVYLA |
|--------------|--|---|
| SEC ID NO:30 | SEQ 1D NO:9 FKSF11VARDSENSLESSHVLEISSGSDIVUSVWGIAKRAGKGVCVLSGTGAVINVILKQ SEO IS NO:10 PKPPIIVTRDSPNALHSHVLEVAGGADVVDCVAEYARRAGRGVCVLSGGGAVVNVALRO | VVILHGRFEILSLIGTSLPPPAPPGAGGLIVILF VATLRGRFEILSLTGTVILPPPAPPGASGLIVFLS |
| SEQ ID NO:11 | SEQ ID NO:11 -K-P-IRDSPN-L-SHVIEG-DV-YA-RRG-GVG-G-V-NVRQV-TI-GRE-IISIIGG-IPPPAPPGA-GIIA | V-TL-CRP-ILSLTGT-LPPPAPPGA-CLL |